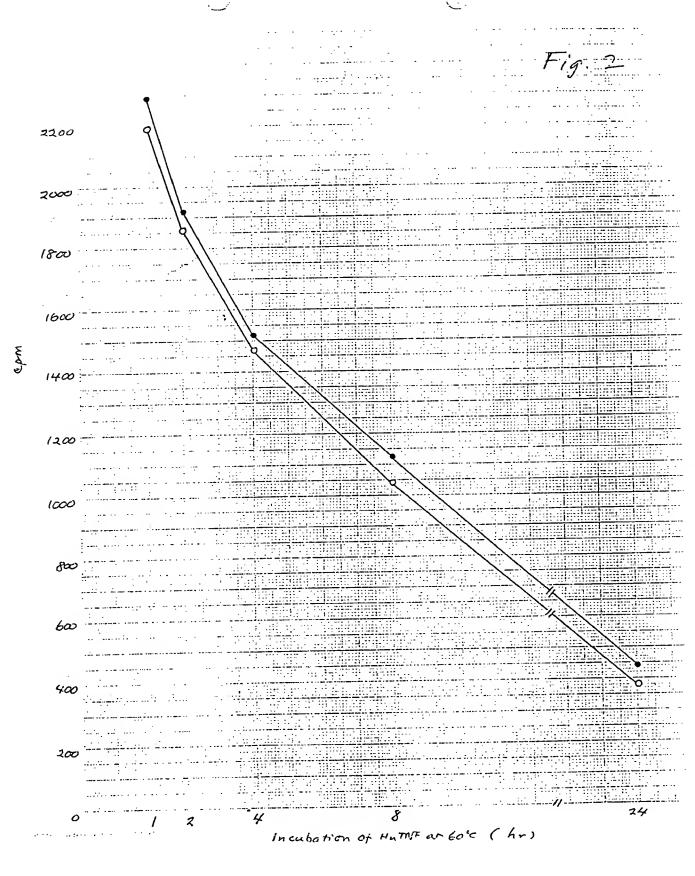
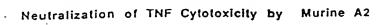


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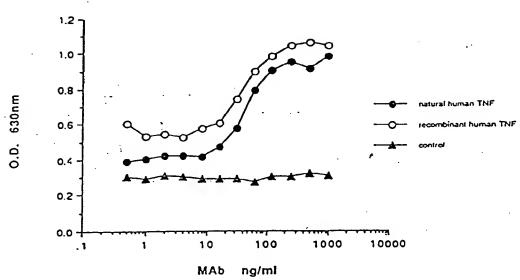
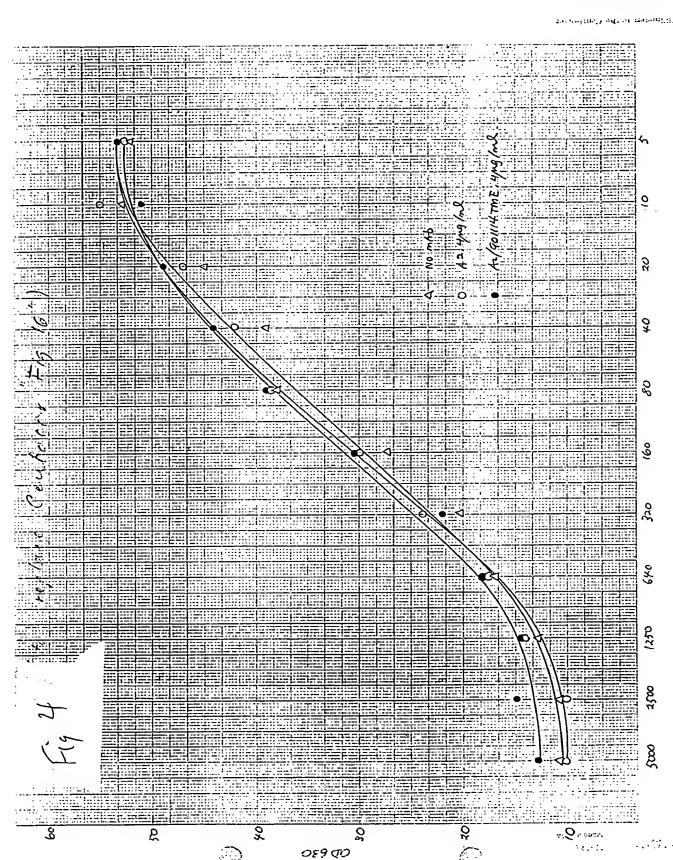
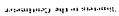


FIG. 3





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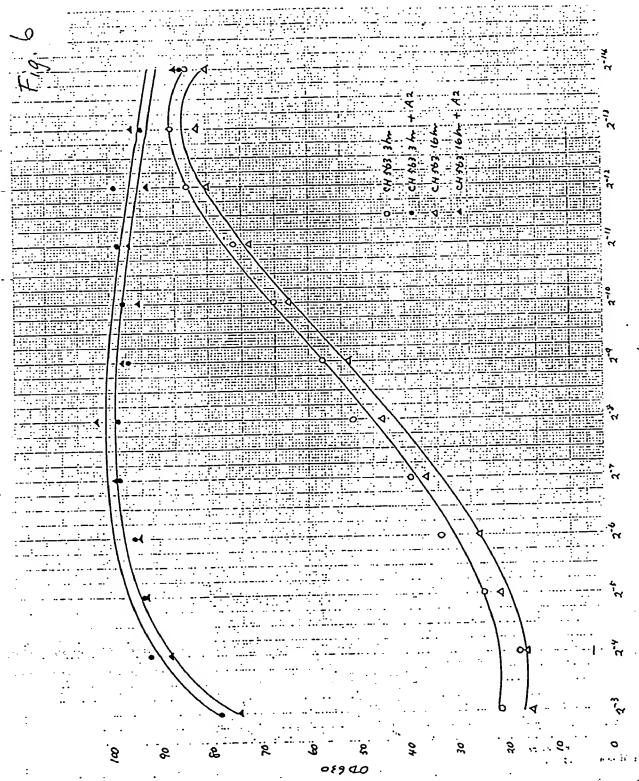
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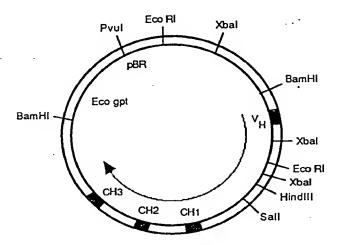


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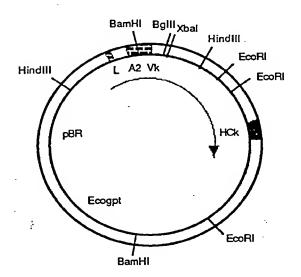
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Fig 8

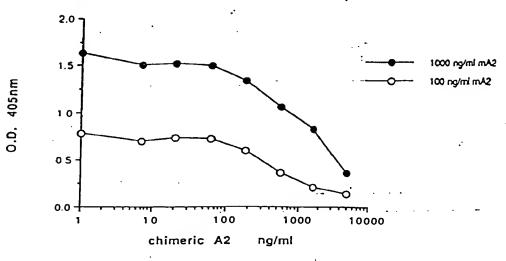


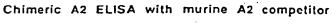
pA2HG1apgpt

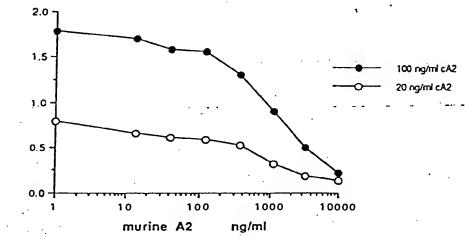


pA2HuKapgpt

Murine A2 ELISA with chimeric A2 competitor

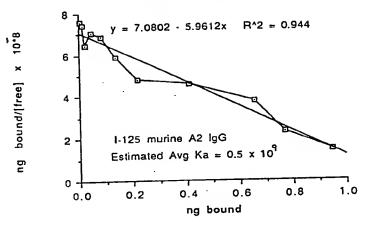






F16. 9

Scatchard Analysis of I-125 mA2 and cA2 IgG



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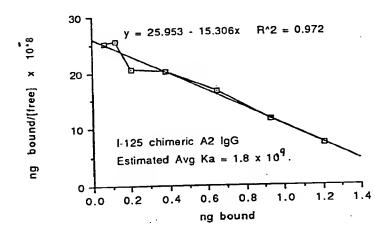
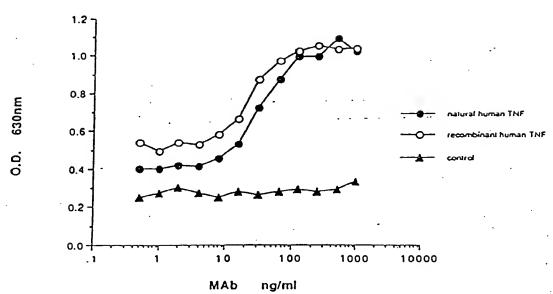


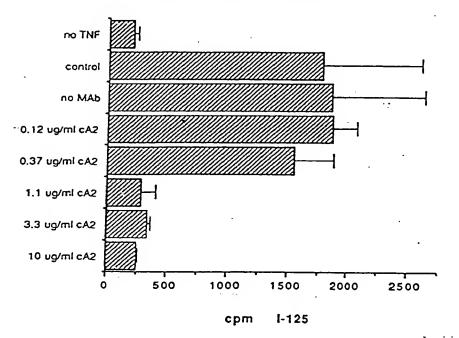
Fig. 10





F16. 11

#### Neutralization of ELAM-1 Expression by Chimeric A2



F10. 12

#### Aur ....

## Figure 13

# SEQUENCE OF HUMAN TNF

10 Ser Arg Thr Pro Ser Asp Lys Pro Val Ala His Val Val Ala Asn Pro Ser Val Arg Ser

Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg Ala Asn Ala Leu Leu Ala Asn Gly 22 Gh ,

Ser Ţ, 50 Glu Leu Arg Asp Asn Gln Leu Val Val Pro Ser Glu Gly Leu Tyr Leu Ile 41 Val

Thr 11e 70 Val Leu Phe Lys Gly Gln Gly Cys Pro Ser Thr His Val Leu Leu Thr His 61 Gh

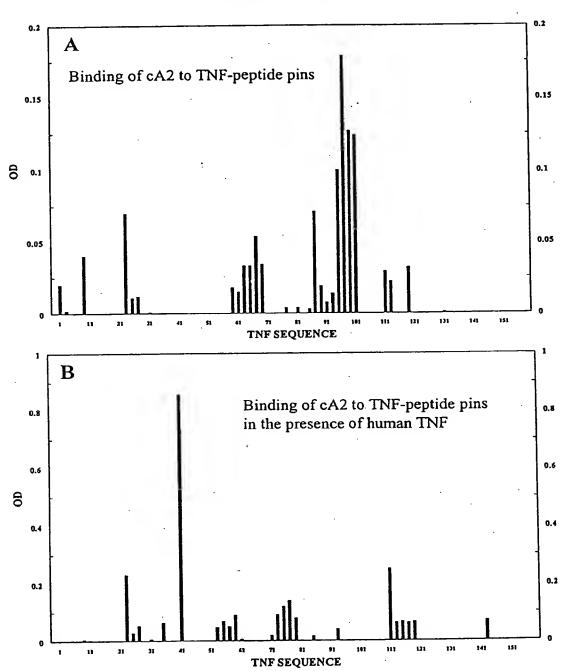
Pro 90 Ser Tyr Gln Thr Lys Val Asn Leu Leu Ser Ala Ile Lys Ser Ala Val 81 Ser Arg Ile

Leu <u>7</u> 101 Cys Gln Arg Glu Thr Pro Glu Gly Ala Glu Ala Lys Pro Trp Tyr Glu Pro Ile

121 Gly Gly Val Phe Gln Leu Glu Lys Gly Asp Arg Leu Ser Ala Glu Ile Asn Arg Pro Asp

Ile Ala Leu 141 Tyr Leu Asp Phe Ala Glu Ser Gly Gln Val Tyr Phe Gly Ile

Figure 14



### Figure 15

# PEPTIDE SEQUENCE OF HUMAN TNF RECOGNIZED BY cA2

1 Val Arg Ser Ser Arg Thr Pro Ser Asp Lys Pro Val Ala His Val Val Ala Asn Pro

21 Gin Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg Ala Asn Ala Leu Leu Ala Asn Gly

41 Val Glu Leu Arg Asp Asn Gln Leu Val Val Pro Ser Glu Gly Leu Tyr Leu Ile

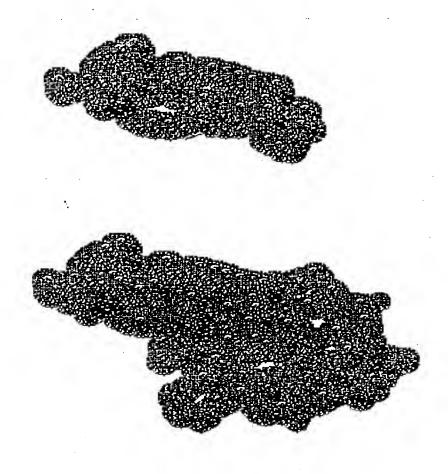
Ser Arg Ile Ala Val Ser Exp. Gin Thr Lys Vall Asin

Tyr Leu 110 Ala Glu Ala Lys Pro Trp Tyr Glu Pro Ile

121 Gly Gly Val Phe Gln Leu Glu Lys Gly Asp Arg Leu Ser Ala Glu Ile Asn Arg Pro Asp

141 Tyr Leu Asp Phe Ala Glu Ser Gly Gln Val Tyr Phe Gly Ile Ile Ala Leu





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#### Figure 17A

#### CA2 LIGHT CHAIN VARIABLE REGION SEQUENCE

GACATCTTCCTGACTCAGTCTCCAGCCATCCTGTCTGTCAGTCCAGGACAAAGAGTCACT AspIleLeuLeuThrGlnSerProAlaIleLeuSerValSerProGlyGluArgValSer

TTCTCCTCCACCCCAGTCAGTTCGTTGGCTCAAGCATCCACTGGTATCAGCAAAGAACA PheSerCysArgAlaSerGlnPheValGlySerSerIleHisTrpTyrGlnGlnArgThr

AATCCTTCTCCAAGGCTTCTCATAAACTATGCTTCTGAGTCTATGTCTGGGATCCCTTCC AsnGlySerProArgLeuLeuIleLysTyrAlaSerGluSerMetSerGlyIleProSer

 $\label{local} \textbf{AGGTTTAGTGGGAGTGGATCAGCGAGAGATTTTAGTGTTAGGATGAAGAGTGTGGAGTGTAGGPheSerGlySerGlySerGlyThrAspPheThrLeuSerIleAsnThrValGluSer$ 

 ${\tt GAAGATATTGCAGATTATTACTGTCAAGAAAGTCATACCTCCCCATTCACCTTCCCCTCC} \\ {\tt GluAspIleAlaAspTyrTyrCysGlnGlnSerHisSerTrpProPheThrPheGlySer} \\$ 

GGGACAAATTTGGAAGTAAAA GlyThrAsnLeuGluValLys

91.23.73 FL 1.23.73  ${\tt CAAGTCAACCTTGACCACTTGCACCACCTTCCTCCAACCTTCCATCAAACTCC1} \\ {\tt CluValLysLeuGluGluSerGlyGlyGlyLeuValGlnProGlyGlySerMetLysLeu} \\$ 

1 9438 T

TCCTGTGTTGCCTCTGGATTCATTTTCAGTAACCACTGGATGAACTGGGTCCGCCAGTCT SerCysValAlaSerClyPheIlePheSerAsnHisTrpMetAsnTrpValArgGlnSer

CCAGAGAACCGGCTTGAGTGGGTTGCTGAAATTAGATCAAAATCTATTAATTCTGCAACA ProGluLysGlyLeuGluTrpValA1aGluI1eArgSerLysSerI1eAsnSerA1aThr

CATTATGCGGAGTCTGTGAAAGGGAGGTTCACCATCTCAAGAGATGATTCCAAAAGTCCT HisTyrAlaGluSerValLysGlyArgPheThrlleSerArgAspAspSerLysSerAla

 ${\tt GTCTAGCTGCAAATGACCGACTTAAGAACTGAAGACACTGGCGTTTATTACTGTTCCAGG} \\ {\tt ValTyrLeuGlnMetThrAspLeuArgThrCluAspThrClyValTyrTyrCysSerArg} \\$ 

Figure 17B

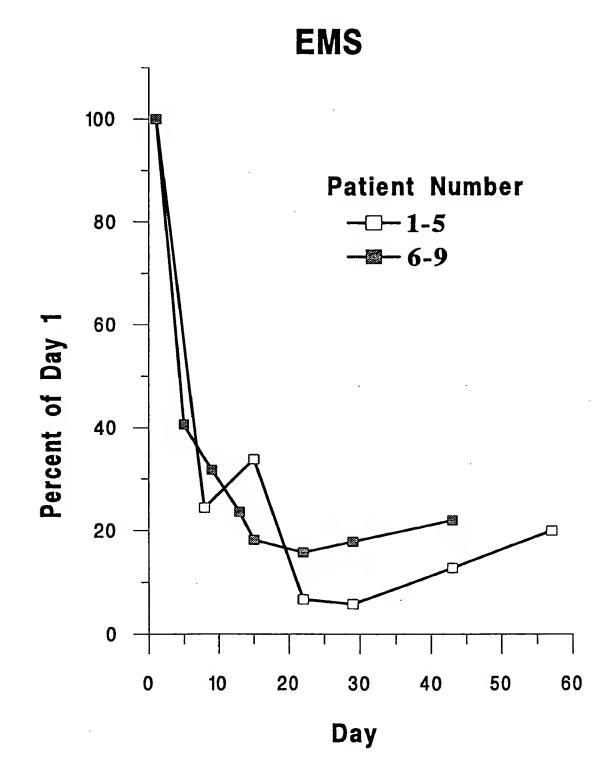
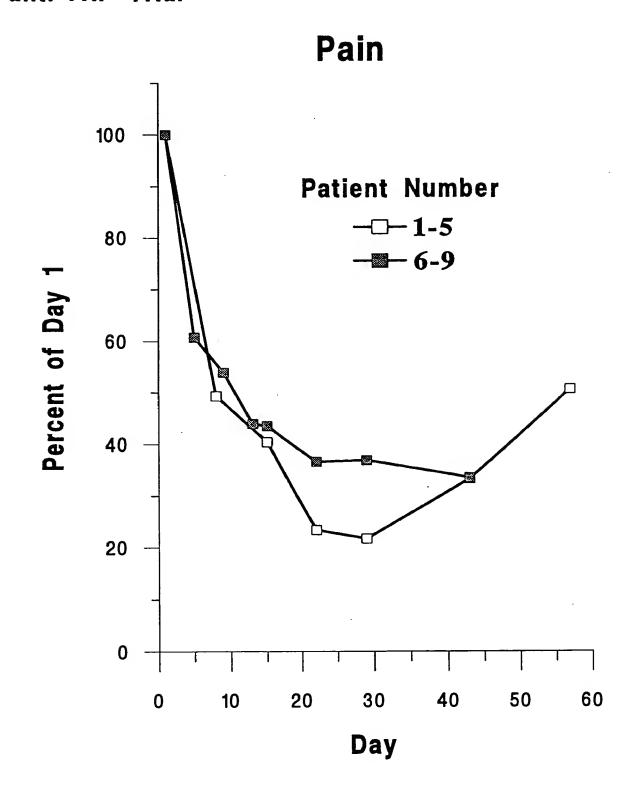
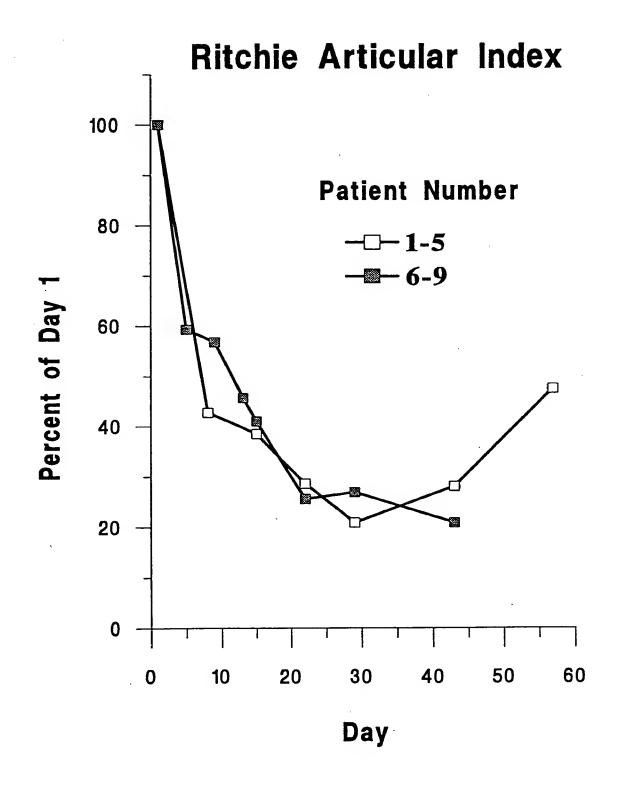


FIG. IS



F16.19

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F16. 20

#### **Swollen Joints**

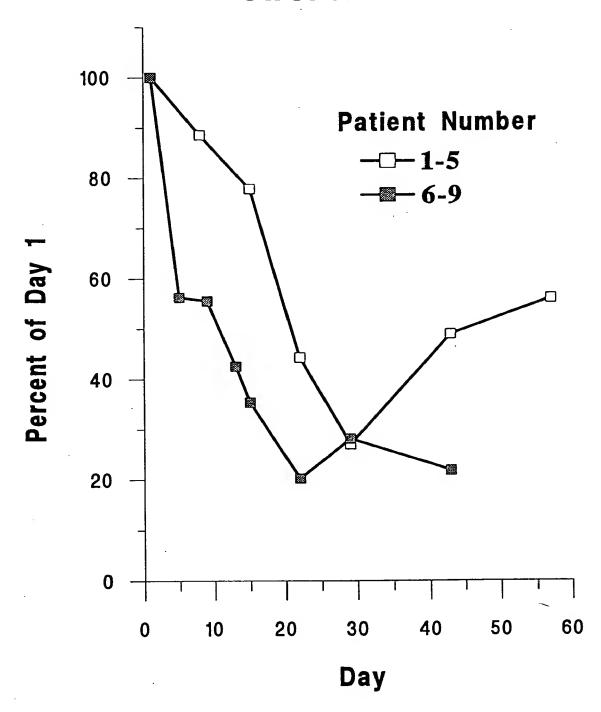
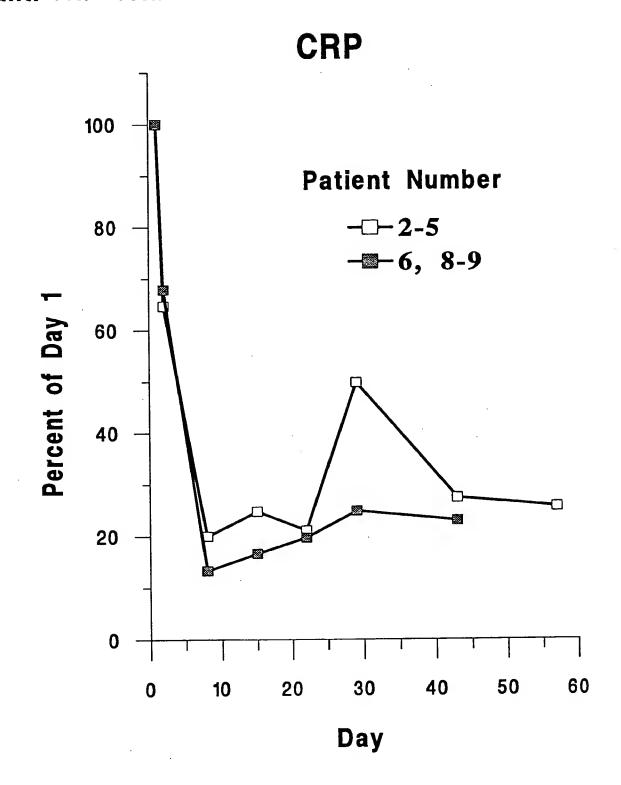


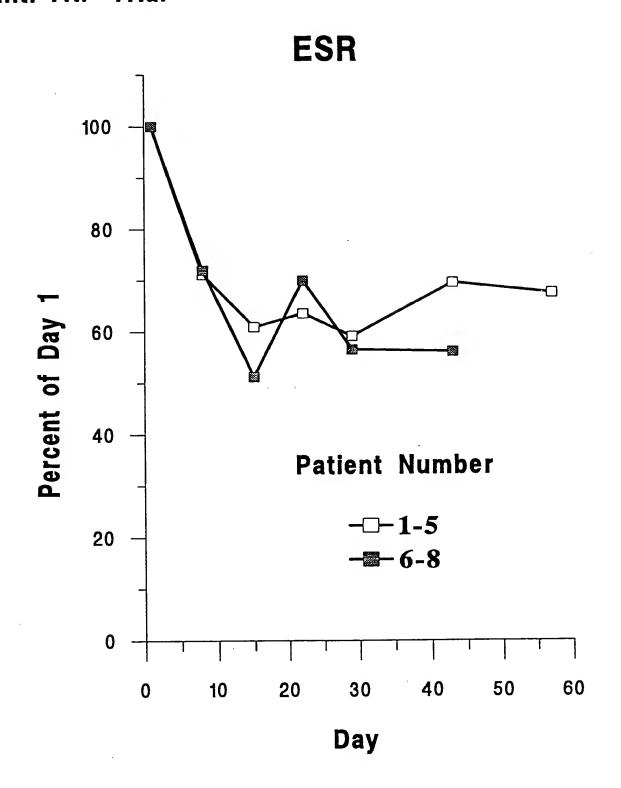
FIG 21

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F16. 22





F16 23



#### **Index of Disease Activity**

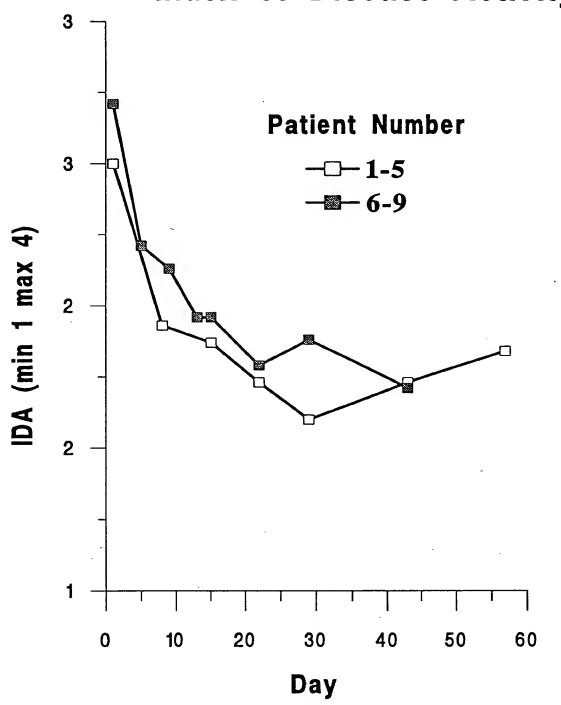


FIG 24